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RAW SEQUENCE LISTING

DATE: 04/25/2002

PATENT APPLICATION: US/10/032,106

TIME: 15:14:47

Input Set : A:\43311-20007.00.txt

Output Set: N:\CRF3\04252002\J032106.raw

ENTERED

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4 <110> APPLICANT: Peng, Zaoyuan
5      Yu, Zailin
6      Wiley, Andrew
7      Hu, Qianjin
9 <120> TITLE OF INVENTION: METHODS FOR IDENTIFYING G-PROTEIN
10     COUPLED RECEPTORS ASSOCIATED WITH DISEASES
13 <130> FILE REFERENCE: 433112000700
15 <140> CURRENT APPLICATION NUMBER: US 10/032,106
C--> 16 <141> CURRENT FILING DATE: 2002-04-08
18 <150> PRIOR APPLICATION NUMBER: US 60/258,070
19 <151> PRIOR FILING DATE: 2000-12-20
21 <160> NUMBER OF SEQ ID NOS: 10
23 <170> SOFTWARE: FastSEQ for Windows Version 4.0
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 924
27 <212> TYPE: DNA
28 <213> ORGANISM: Homo sapiens
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32 tggcaagttc tcttcttcat catcttttctt gtggtctaca tcatacccat ggtgggcaat      120
33 atcggcatga tgggtgtaat caaggtcagt cctcagctta acaaccccat gtactttttc      180
34 ctcatgact tgtcatttgt tgatgtgtgg ttttcttcca atgtcacccc taaaatgttg      240
35 gaaaacctgt tttcagataa aaaaacaatt acttatgctg gttgtttagt acagtgtttc      300
36 ttcttcattg ctcttgtcca tgtggaaatt tttattcttg ctgcgatggc ctttgataga      360
37 tacatggcaa ttgggaatcc tctgctttat ggcagtaaaa tgtcaagggg tgtctgtatt      420
38 cgactgatta ctttccctta catttatggg tttctgacga gtctggcagc aacattatgg      480
39 acttacggct tgtacttctg tggaaaaatt gagatcaacc atttctactg tgcagatcca      540
40 cctctcatca aaatggcctg tgccgggacc tttgtaaaag aatatacaat gatcatactt      600
41 gccggcatta acttcacata ttccctgact gtaattatca tctcttactt attcactctc      660
42 attgccattc tgcgaatgcg ctacagcagaa ggaaggcaga aggccttttc cacatgtggg      720
43 tcccatctga cagctgtcat tatattctat ggtactctga tcttcatgta tctcagacgt      780
44 cccacagagg agtctgtgga gcagggggaa atggtggctg tgttctatac cacagtgatc      840
45 cccatgttga atcccatgat ctacagtctg aggaacaagg atgtgaaaaa ggccatgatg      900
46 aaagtgatca gcagatcatg ttaa                                     924
48 <210> SEQ ID NO: 2
49 <211> LENGTH: 939
50 <212> TYPE: DNA
51 <213> ORGANISM: Homo sapiens
53 <400> SEQUENCE: 2
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55 cctcagctgg agctagtcct ctttgtgggt cttttgatct tctatatctt cactttgctg      120
56 gggaaacaaa ccatcattgt attatctcac ttggaccac atcttcacac tctatgtat      180
57 tttttcttct ccaacctaa gctttttggat ctgtgttaca caaccggcat tgttccacag      240

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58 ctccctgggta atctcagggg agcagacaaa tcaatctcct atgggtgggtg tgtagttcag 300
59 ctgtacatct ctctaggctt gggatctaca gaatgcgttc tcttaggagt gatggatattt 360
60 gaccgctatg cagctgtttg caggccccctc cactacacag tagtcatgca cccttgtctg 420
61 tatgtgctga tggcttctac ttcattgggtc attgggtttg ccaactccct attgcagacg 480
62 gtgctcatct tgcttttaac actttgtgga agaaataaat tagaacaact tctttgtgag 540
63 gttcctccat tgetcaagct tgccgtgtgt gacactacta tgaatgaatc tgaactcttc 600
64 tttgtcagtg tcattattct tctgtacct gttgcattaa tcatattctc ctatagtcag 660
65 attgtcaggg cagtcacag gataaagtta gcaacagggc agagaaaagt gtttgggaca 720
66 tgtggctccc acctcacagt ggtttccctg ttctacggca cagctatcta tgcttacctc 780
67 cagcccgga acaactactc tcaggatcag ggcaagttca tctctctctt ctacaccatc 840
68 attacacca tgatcaaccc cctcatatat acactgagga acaaggatgt gaaaggagca 900
69 cttaagaagg tgctctggaa gaactacgac tccagatga 939

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71 <210> SEQ ID NO: 3

72 <211> LENGTH: 930

73 <212> TYPE: DNA

74 <213> ORGANISM: Homo sapiens

76 <220> FEATURE:

77 <221> NAME/KEY: misc_feature

78 <222> LOCATION: (1)...(930)

79 <223> OTHER INFORMATION: n= a, c, g, or t

81 <400> SEQUENCE: 3

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82 atggctgcag gaaatcactc tacagtgcag gagttcattc tcaaggggtt aacgaagaga 60
83 gcagacctcc agctccccct ctttctcttc ttctcggga tctacttggt caccatcgtg 120
84 gggaaacctg gcatgatcac tctaatttgt ctgaactctc agctgcacac ccccatgtac 180
85 tactttctca gcaatctgtc actcatggat ctctgctact cctcogtcat taccctaag 240
86 atgctgggta actttgtgac agagaaaaaac atcatctcct acgcaggggtg catgtcacag 300
87 ctctacttct tctttgtttt tgtcattgct gagtggtaca tgetgacagt gatggcctac 360
W--> 88 gaccgctatg ttgnctctg ccaccctttg ctttacaaca tcattatgtc tcatcacacc 420
89 tgctgtctgc tgggtgctgt ggtctacgcc atcggactca ttggctccac aatagaaact 480
90 ggcctcatgt taaaactgcc ctattgtgag caccctcatc gtcactactt ctgtgacatc 540
91 ctccctctca tgaagctgtc ctgctctagc acctatgatg ttgagatgac agtcttcttt 600
92 tcggctggat tcaacatcat agtcacgagc ttaacagttc ttgtttctta caccttcatt 660
93 ctctccagca tctcggcat cagcaccaca gaggggagat ccaaagcctt cagcacctgc 720
94 agtccccacc ttgcagccgt gggaatgttc tatggatcaa ctgcattcat gtacttaaaa 780
95 cctccacaaa tcagttcctt gaccacaggag aatgtggcct ctgtgttcta caccacggtg 840
96 atccccatgt tgaatccctt aatctacagc ctgaggaaca aggaagtaaa ggctgccgtg 900
97 cagaaaacgc tgaggggtaa actgttttga 930

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99 <210> SEQ ID NO: 4

100 <211> LENGTH: 927

101 <212> TYPE: DNA

102 <213> ORGANISM: Homo sapiens

104 <400> SEQUENCE: 4

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105 atggccatgg acaatgtcac agcagtgttt cagtttctcc ttattggcat ttctaactat 60
106 cctcaatgga gagacacgtt ttccacatta gtgctgataa ttacctcag cacattgttg 120
107 gggaatggat ttatgatctt tcttatcacc tttagaccca acctccacac tccaatctac 180
108 ttcttctcta gtaacctgtc ttcttagac ctttggtatg gaacagcttc catgccccag 240
109 gctttgggtg attgtttctc tacccatccc tacctctctt atccccgatg ttggetcaa 300
110 acgagtgtct ccttggcttt ggccacagca gagtgccctc tactggetgc catggcctat 360
111 gaccgtgtgg ttgctatcag caatccccctg cgttattcag tggttatgaa tggcccagta 420

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112 tgtgtctgct tgggtgctac ctcatggggg acatcacttg tgctcactgc catgctcactc 480
113 ctatccctga ggcttcactt ctgtggggct aatgtcatca accattttgc ctgtgagatt 540
114 ctctccctca ttaagctgac ctgttctgat accagcctca atgaatttat gatcctcactc 600
115 accagtatct tcaccctgct gctaccattt gggtttgctc tcctctccta catacgaatt 660
116 gctatggcta tcataaggat tcgctcactc cagggcaggc tcaaggcctt taccacatgt 720
117 ggctctcacc tgaccgtggg gacaatcttc tatgggtcag ccactctccat gtatatgaaa 780
118 actcagtcga agtccctacc tgaccaggac aagtttatct cagtgtttta tggagctttg 840
119 acacccatgt tgaacccctt gatatatagc ctgagaaaaa aagatgttaa acgggcaata 900
120 aggaaagtta tgttgaaaag gacatga 927

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123 <211> LENGTH: 945

124 <212> TYPE: DNA

125 <213> ORGANISM: Homo sapiens

127 <400> SEQUENCE: 5

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129 ggcttggaac acctgcacat ctggatctcc atccctttct gcttagcata tacactggcc 120
130 ctgcttgga actgcactct ccttctcact atccaggctg atgcagcctt ccatgaaccc 180
131 atgtacctct ttctggccat gttggcagcc atcgacctgg tcctttcctc ctgagcactg 240
132 cccaaaatgc ttgccatatt ctggttcagg gatcgggaga taaacttctt tgctgtctg 300
133 gccagatgt tcttccttca ctcttctctc atcatggagt cagcagtgtt gctggccatg 360
134 gcctttgacc gctatgtggc tatctgcaag ccaactgact acaccaaggc cctgactggg 420
135 tccctcatca ccaagattgg catggctgct gtggcccggg ctgtgacact aatgactcca 480
136 ctcccttcc tgctgagatg ttccactac tgccgaggcc cagtgatcgc tcaactgtac 540
137 tgtgaacaca tggctgtggt gaggtggtg tgtggggaca ctgcttcaa caatatctat 600
138 ggcatcgctg tggccatggt tattgtggtg ttggacctgc tccttggtat cctgtcttat 660
139 atctttattc ttcaggcagt tctactgctt gcctctcagg aggcccgcta caaggcattt 720
140 gggacatgtg tctctcatat aggtgccatc ttagccttct acacaactgt ggtcatctct 780
141 tcagtcatgc accgtgtagc ccgccatgct gccctcatg tccacatcct ccttgccaat 840
142 ttctatctgc tcttcccacc catggtcaat cccataatct atggtgtcaa gaccaagcaa 900
143 atccgtgaga gcatcttggg agtattccca agaaaggata tgtag 945

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145 <210> SEQ ID NO: 6

146 <211> LENGTH: 307

147 <212> TYPE: PRT

148 <213> ORGANISM: Homo sapiens

150 <400> SEQUENCE: 6

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152 1 5 10 15
153 Ser Arg Arg Glu Trp Gln Val Leu Phe Phe Ile Ile Phe Leu Val Val
154 20 25 30
155 Tyr Ile Ile Thr Met Val Gly Asn Ile Gly Met Met Val Leu Ile Lys
156 35 40 45
157 Val Ser Pro Gln Leu Asn Asn Pro Met Tyr Phe Phe Leu Ser His Leu
158 50 55 60
159 Ser Phe Val Asp Val Trp Phe Ser Ser Asn Val Thr Pro Lys Met Leu
160 65 70 75 80
161 Glu Asn Leu Phe Ser Asp Lys Lys Thr Ile Thr Tyr Ala Gly Cys Leu
162 85 90 95
163 Val Gln Cys Phe Phe Phe Ile Ala Leu Val His Val Glu Ile Phe Ile
164 100 105 110

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165 Leu Ala Ala Met Ala Phe Asp Arg Tyr Met Ala Ile Gly Asn Pro Leu
166      115      120      125
167 Leu Tyr Gly Ser Lys Met Ser Arg Val Val Cys Ile Arg Leu Ile Thr
168      130      135      140
169 Phe Pro Tyr Ile Tyr Gly Phe Leu Thr Ser Leu Ala Ala Thr Leu Trp
170 145      150      155      160
171 Thr Tyr Gly Leu Tyr Phe Cys Gly Lys Ile Glu Ile Asn His Phe Tyr
172      165      170      175
173 Cys Ala Asp Pro Pro Leu Ile Lys Met Ala Cys Ala Gly Thr Phe Val
174      180      185      190
175 Lys Glu Tyr Thr Met Ile Ile Leu Ala Gly Ile Asn Phe Thr Tyr Ser
176      195      200      205
177 Leu Thr Val Ile Ile Ile Ser Tyr Leu Phe Ile Leu Ile Ala Ile Leu
178      210      215      220
179 Arg Met Arg Ser Ala Glu Gly Arg Gln Lys Ala Phe Ser Thr Cys Gly
180 225      230      235      240
181 Ser His Leu Thr Ala Val Ile Ile Phe Tyr Gly Thr Leu Ile Phe Met
182      245      250      255
183 Tyr Leu Arg Arg Pro Thr Glu Glu Ser Val Glu Gln Gly Lys Met Val
184      260      265      270
185 Ala Val Phe Tyr Thr Thr Val Ile Pro Met Leu Asn Pro Met Ile Tyr
186      275      280      285
187 Ser Leu Arg Asn Lys Asp Val Lys Lys Ala Met Met Lys Val Ile Ser
188      290      295      300
189 Arg Ser Cys
190 305
192 <210> SEQ ID NO: 7
193 <211> LENGTH: 312
194 <212> TYPE: PRT
195 <213> ORGANISM: Homo sapiens
197 <400> SEQUENCE: 7
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199 1      5      10      15
200 Phe Ser Asp Arg Pro Gln Leu Glu Leu Val Leu Phe Val Val Leu Leu
201      20      25      30
202 Ile Phe Tyr Ile Phe Thr Leu Leu Gly Asn Lys Thr Ile Ile Val Leu
203      35      40      45
204 Ser His Leu Asp Pro His Leu His Thr Pro Met Tyr Phe Phe Phe Ser
205      50      55      60
206 Asn Leu Ser Phe Leu Asp Leu Cys Tyr Thr Thr Gly Ile Val Pro Gln
207 65      70      75      80
208 Leu Leu Val Asn Leu Arg Gly Ala Asp Lys Ser Ile Ser Tyr Gly Gly
209      85      90      95
210 Cys Val Val Gln Leu Tyr Ile Ser Leu Gly Leu Gly Ser Thr Glu Cys
211      100      105      110
212 Val Leu Leu Gly Val Met Val Phe Asp Arg Tyr Ala Ala Val Cys Arg
213      115      120      125
214 Pro Leu His Tyr Thr Val Val Met His Pro Cys Leu Tyr Val Leu Met
215      130      135      140

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216 Ala Ser Thr Ser Trp Val Ile Gly Phe Ala Asn Ser Leu Leu Gln Thr
217 145 150 155 160
218 Val Leu Ile Leu Leu Leu Thr Leu Cys Gly Arg Asn Lys Leu Glu His
219 165 170 175
220 Phe Leu Cys Glu Val Pro Pro Leu Leu Lys Leu Ala Cys Val Asp Thr
221 180 185 190
222 Thr Met Asn Glu Ser Glu Leu Phe Phe Val Ser Val Ile Ile Leu Leu
223 195 200 205
224 Val Pro Val Ala Leu Ile Ile Phe Ser Tyr Ser Gln Ile Val Arg Ala
225 210 215 220
226 Val Met Arg Ile Lys Leu Ala Thr Gly Gln Arg Lys Val Phe Gly Thr
227 225 230 235 240
228 Cys Gly Ser His Leu Thr Val Val Ser Leu Phe Tyr Gly Thr Ala Ile
229 245 250 255
230 Tyr Ala Tyr Leu Gln Pro Gly Asn Asn Tyr Ser Gln Asp Gln Gly Lys
231 260 265 270
232 Phe Ile Ser Leu Phe Tyr Thr Ile Ile Thr Pro Met Ile Asn Pro Leu
233 275 280 285
234 Ile Tyr Thr Leu Arg Asn Lys Asp Val Lys Gly Ala Leu Lys Lys Val
235 290 295 300
236 Leu Trp Lys Asn Tyr Asp Ser Arg
237 305 310
239 <210> SEQ ID NO: 8
240 <211> LENGTH: 309
241 <212> TYPE: PRT
242 <213> ORGANISM: Homo sapiens
244 <220> FEATURE:
245 <221> NAME/KEY: VARIANT
246 <222> LOCATION: (1)...(309)
247 <223> OTHER INFORMATION: Xaa= any amino acid
249 <400> SEQUENCE: 8
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252 Leu Thr Lys Arg Ala Asp Leu Gln Leu Pro Leu Phe Leu Leu Leu
253 20 25 30
254 Gly Ile Tyr Leu Val Thr Ile Val Gly Asn Leu Gly Met Ile Thr Leu
255 35 40 45
256 Ile Cys Leu Asn Ser Gln Leu His Thr Pro Met Tyr Tyr Phe Leu Ser
257 50 55 60
258 Asn Leu Ser Leu Met Asp Leu Cys Tyr Ser Ser Val Ile Thr Pro Lys
259 65 70 75 80
260 Met Leu Val Asn Phe Val Ser Glu Lys Asn Ile Ile Ser Tyr Ala Gly
261 85 90 95
262 Cys Met Ser Gln Leu Tyr Phe Phe Leu Val Phe Val Ile Ala Glu Cys
263 100 105 110
W--> 264 Tyr Met Leu Thr Val Met Ala Tyr Asp Arg Tyr Val Xaa Xaa Cys His
265 115 120 125
266 Pro Leu Leu Tyr Asn Ile Ile Met Ser His His Thr Cys Leu Leu Leu
267 130 135 140

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/032,106

DATE: 04/25/2002
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Input Set : A:\43311-20007.00.txt
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; N Pos. 374,376
Seq#:8; Xaa Pos. 125,126